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Input Set : A:\EP.txt

Output Set: N:\CRF3\06032002\J014320.raw

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3 <110> APPLICANT: North Shore - Long Island Jewish Research Institute
         Tabibzadeh, Siamak
 6 <120> TITLE OF INVENTION: METHOD FOR DIAGNOSING A PRE-NEOPLASTIC OR NEOPLASTIC LESION
         TRANSITIONAL EPITHELIAL CELLS
9 <130> FILE REFERENCE: 50425/137
11 <140> CURRENT APPLICATION NUMBER: 10/014,320
12 <141> CURRENT FILING DATE: 2001-12-11
14 <150> PRIOR APPLICATION NUMBER: 60/255,641
15 <151> PRIOR FILING DATE: 2000-12-14
17 <160> NUMBER OF SEQ ID NOS: 6
19 <170> SOFTWARE: PatentIn version 3.1
21 <210> SEO ID NO: 1
22 <211> LENGTH: 1161
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
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                                                                         120
31 gagcagetee tgggcageet getgeggcag etgeagetea gegaggtgee egtaetggae
                                                                         180
33 agggccgaca tggagaagct ggtcatcccc gcccacgtga gggcccagta tgtagtcctg
                                                                         240
35 ctgcggcgca gccacgggga ccgctcccgc ggaaagaggt tcagccagag cttccgagag
                                                                         300
37 gtggccggca ggttcctggc gtcggaggcc agcacacacc tgctggtgtt cggcatggag
                                                                         360
39 cageggetge egeceaacag egagetggtg caggeegtge tgeggetett eeaggageeg
                                                                         420
41 gtccccaagg ccgcgctgca caggcacggg cggctgtccc cgcgcagcgc ccaggcccgg
                                                                         480
43 gtgaccgtcg agtggctgcg cgtccgcgac gacggctcca accgcacctc cctcatcgac
                                                                         540
45 tecaggetgg tgteegteea egagagegge tggaaggeet tegaegtgae egaggeegtg
                                                                         600
47 aacttetgge ageagetgag eeggeeeegg eageegetge tgetaeaggt gteggtgeag
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49 agggagcate tgggcccgct ggcgtccggc gcccacaagc tggtccgctt tgcctcgcag
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51 ggggcgccag ccgggcttgg ggagccccag ctggagctgc acaccctgga cctcagggac
                                                                         780
53 tatggagete agggegaetg tgaecetgaa geaceaatga eegagggeae eegetgetge
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55 cgccaggaga tgtacattga cctgcagggg atgaagtggg ccaagaactg ggtgctggag
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57 cccccgggct tcctggctta cgagtgtgtg ggcacctgcc agcagccccc ggaggccctg
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59 gccttcaatt ggccatttct ggggccgcga cagtgtatcg cctcggagac tgcctcgctg
                                                                        1020
61 cccatgatcg tcagcatcaa ggagggaggc aggaccaggc cccaggtggt caqcctgccc
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63 aacatgaggg tgcagaagtg cagctgtgcc tcggatgggg cgctcgtgcc aaggaggctc
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68 <210> SEQ ID NO: 2
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71 <213> ORGANISM: Homo sapiens
73 <400> SEQUENCE: 2
75 Met Trp Pro Leu Trp Leu Cys Trp Ala Leu Trp Val Leu Pro Leu Ala
76 1
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79 Gly Pro Gly Ala Ala Leu Thr Glu Glu Gln Leu Leu Gly Ser Leu Leu
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83 Arg Gln Leu Gln Leu Ser Glu Val Pro Val Leu Asp Arg Ala Asp Met
                              40
87 Glu Lys Leu Val Ile Pro Ala His Val Arg Ala Gln Tyr Val Val Leu
                          55
91 Leu Arg Arg Ser His Gly Asp Arg Ser Arg Gly Lys Arg Phe Ser Gln
                      70
                                          75
95 Ser Phe Arg Glu Val Ala Gly Arg Phe Leu Ala Ser Glu Ala Ser Thr
99 His Leu Leu Val Phe Gly Met Glu Gln Arg Leu Pro Pro Asn Ser Glu
               100
                                   105
103 Leu Val Gln Ala Val Leu Arg Leu Phe Gln Glu Pro Val Pro Lys Ala
104 115
                               120
107 Ala Leu His Arg His Gly Arg Leu Ser Pro Arg Ser Ala Gln Ala Arg
                           135
                                               140
111 Val Thr Val Glu Trp Leu Arg Val Arg Asp Asp Gly Ser Asn Arg Thr
                       150
                                           155
115 Ser Leu Ile Asp Ser Arg Leu Val Ser Val His Glu Ser Gly Trp Lys
                   165
                                       170
119 Ala Phe Asp Val Thr Glu Ala Val Asn Phe Trp Gln Gln Leu Ser Arg
               180
                                   185
123 Pro Arg Gln Pro Leu Leu Gln Val Ser Val Gln Arg Glu His Leu
           195
                               200
127 Gly Pro Leu Ala Ser Gly Ala His Lys Leu Val Arg Phe Ala Ser Gln
                           215
131 Gly Ala Pro Ala Gly Leu Gly Glu Pro Gln Leu Glu Leu His Thr Leu
                       230
                                           235
135 Asp Leu Arg Asp Tyr Gly Ala Gln Gly Asp Cys Asp Pro Glu Ala Pro
                   245
                                       250
139 Met Thr Glu Gly Thr Arg Cys Cys Arg Gln Glu Met Tyr Ile Asp Leu
               260
                                   265
143 Gln Gly Met Lys Trp Ala Lys Asn Trp Val Leu Glu Pro Pro Gly Phe
147 Leu Ala Tyr Glu Cys Val Gly Thr Cys Gln Gln Pro Pro Glu Ala Leu
                           295
151 Ala Phe Asn Trp Pro Phe Leu Gly Pro Arg Gln Cys Ile Ala Ser Glu
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                                           315
155 Thr Ala Ser Leu Pro Met Ile Val Ser Ile Lys Glu Gly Gly Arg Thr
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                                       330
159 Arg Pro Gln Val Val Ser Leu Pro Asn Met Arg Val Gln Lys Cys Ser
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169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
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	<223> OTHER INFORMATION: forward primer	
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181	<212> TYPE: DNA	
182	<213> ORGANISM: Artificial Sequence	
184	<220> FEATURE:	
185	<223> OTHER INFORMATION: reverse primer	
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205	<212> TYPE: DNA	
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208	<220> FEATURE:	
209	<223> OTHER INFORMATION: reverse primer	
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VERIFICATION SUMMARY DATE: 06/03/2002

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